

RESULT 10
A14250
LOCUS A14250 1021 bp DNA linear PAT 28-JAN-1994
DEFINITION beta-PPT DNA.
ACCESSION A14250
VERSION A14250.1 GI:490128
KEYWORDS .
SOURCE Homo sapiens (human)
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
Hominidae; Homo.
REFERENCE 1 (bases 1 to 1021)
AUTHORS .
TITLE HUMAN TACHYKININS AND THEIR PRECURSORS
JOURNAL Patent: WO 8707643-A 2 17-DEC-1987;
FEATURES Location/Qualifiers
source 1. .1021
/organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"

ORIGIN

Query Match 78.8%; Score 790.4; DB 2; Length 1021;
Best Local Similarity 90.1%; Pred. No. 3.4e-245;
Matches 921; Conservative 0; Mismatches 1; Indels 100; Gaps 3;

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Qy      64 GAGAGTGC GGAGCGAC CAGCGT GCGCTCG GAGGAAC CAGAGAA CT CAGCACCC CGCGGG 123
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Db      1  GAGAGTGC GGAGCGACCA-CGTGCGCTCG GAGGAAC CAGAGAA CT CAGCACCC CGCGGG 59

Qy     124 ACTGTCCGTCGCAA AATCCAACATGAAA ATCTCGTGGCCTTGGCAGTCTTTTCTTGT 183
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Qy     184 CTCCACTCAGCTGTTTGCAGAAGAAATAGGAGCCAATGATGATCTGAATTACTGGTCCGA 243
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Db     120 CTCCACTCAGCTGTTTGCAGAAGAAATAGGAGCCAATGATGATCTGAATTACTGGTCCGA 179

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Db     180 CTGGTACGACAGCGAC CAGATCAAGGAGGA ACTGCCG GAGCCCTTTGAGCATCTTCTGCA 239

Qy     304 GAGAATCGCCCGGAGACCCAAGCCTCAGCAGTTCTTTGGATTAATGGGCAAACGGGATGC 363
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Db     240 GAGAATCGCCCGGAGACCCAAGCCTCAGCAGTTCTTTGGATTAATGGGCAAACGGGATGC 299

Qy     364 -----TGGACATGGCCAGAT 378
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Db     300 TGATTCTCAATTGAAAAACAAGTGGCCCTGTTAAAGGCTCTTTATGGACATGGCCAGAT 359

Qy     379 CTCTCACAAAA----- 389
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Db     360 CTCTCACAAAAGACATAAAACAGATTCCCTTTGTTGGACTAATGGGCAAAGAGCTTTAAA 419

Qy     390 -----TGGCTTATGAAAGGAGTGCAATGCAGAATTATGAAAGAAGACGTTAATAAACTAC 444
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Qy     445 CTAACATTATTTATTTCAGCTTCATTTGTGTCAATGGGCAATGACAGGTAAATTAAGACAT 504
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Db	480	CTAACATTATTTATTCAGCTTCATTTGTGTCAATGGGCAATGACAGGTAAATTAAGACAT	539
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Qy	565	TGTTTATTTTTCATATTGTGCCAATATGTATTGTAAACATGTGTTTTAATTCCAATATGA	624
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Qy	625	TGACTCCCTTAAAATAGAAATAAGTGGTTATTTCTCAACAAAGCACAGTGTTAAATGAAA	684
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Qy	685	TTGTAAACCTGTCAATGATACAGTCCCTAAAGAAAAAAATCATTGCTTTGAAGCAGTT	744
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Qy	805	TTTCATGGTGAAAATGTACTGAGATTTTGGTATTACACTGTATTTGTATCTCTGAAGCAT	864
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Qy	925	TTCATTTTCATTGTATGATGTGTTGTGATAGCTAACATTTTAAATAAAAGAAAAAATATCT	984
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Qy	985	TG 986	
Db	1020	TG 1021	